

FIG. 1

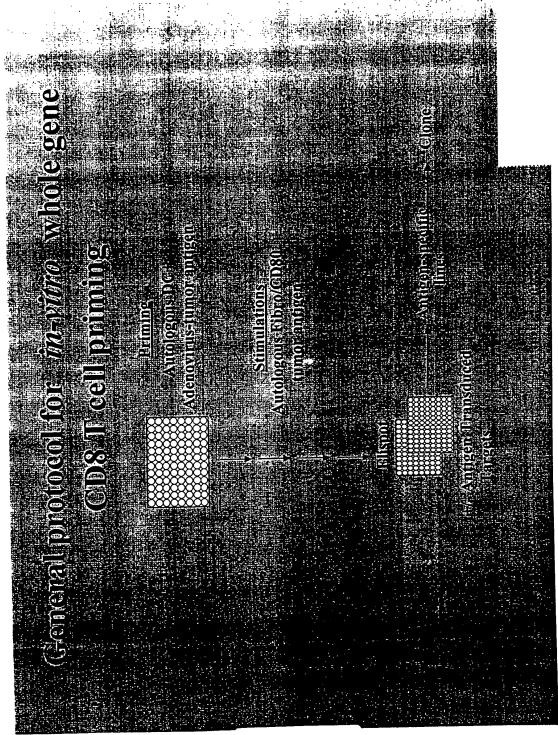


FIG. 2

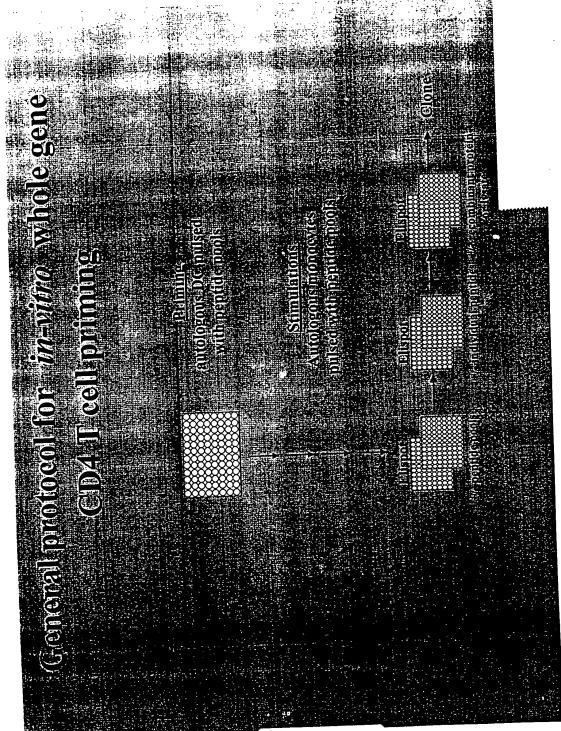


FIG. 3

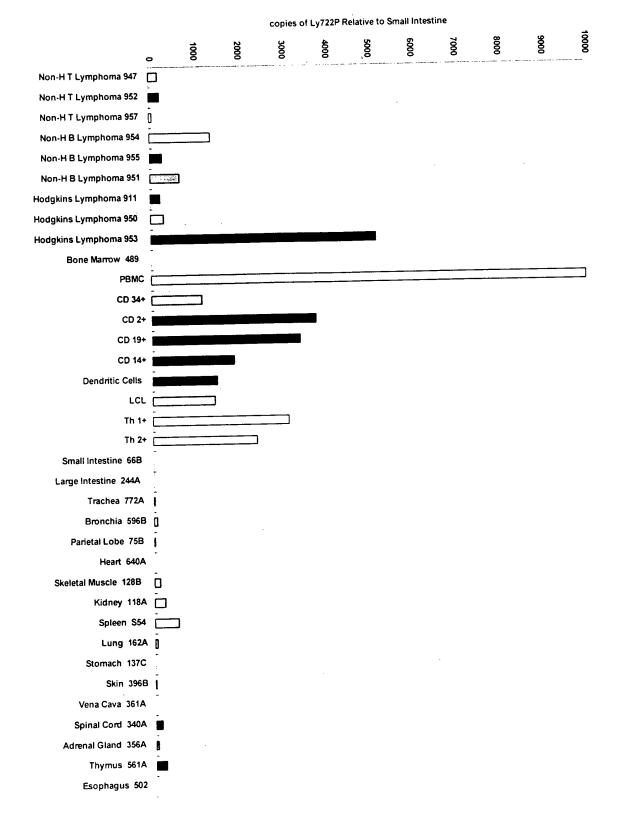


FIG. 4

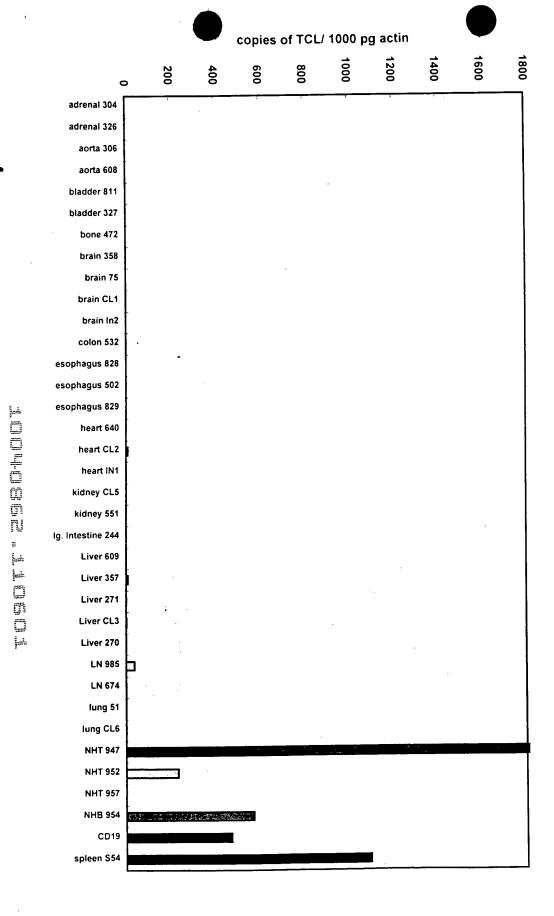
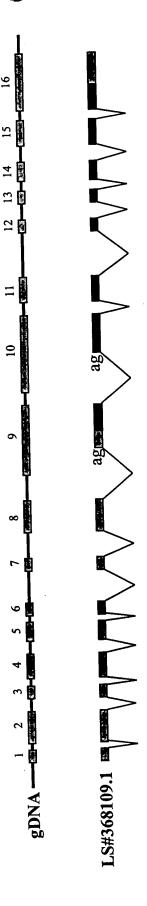


FIG. 5

Chromosome 1:



LS#1384258.1

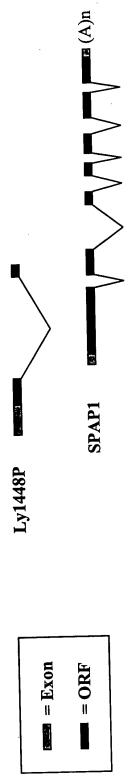


FIG. 6

A - AMPHI mid points of blocks.
R - Residues matching the Rothbard/Taylor motif.
D - Residues matching the IAd motif.
d - Residues matching the IEd motif.

5 MGKKTQR	SLSAE	LEIPA	VKESI AAAA!	DAGKY A	YCRADI	NGHVP.	RRI	RR		וחמת			RRRR.	K
80 ALRGSPI	85 PILYQF	90	95 TLGN:	100 SSAPS	105 GGGAS	110 FNLSL	115 TAEHS	120 GNYSC	125 EANNG	130 LGAQC AAAAA	135 SEAVP A.AAA	140 VSISGI	145 PDGYRF	150 RDLM A .RRR
RRR		RI	KKKK.			· · · · <u>·</u>		• • • •						
155 TAGVLW	160 GLFGVI AAAAA <i>I</i> RRRRRI	165 GFTG' AA RRRRR	170 VALLL	175 YALFH AAAAA	180 KISGE AA	185 SSATN	190 EPRGA AAAAA	195 SRPNP	200 QEFTY .RRRR	205 SSPTP 	210 DMEEL	215 QPVYV	220 NVGSVI	DVDV RR .DDD
230 VYSQVW	235 ISMQQP	240 ESSAN	245 IRTLI AAA .RRRI	250 LENKDS	255 SQVIYS	260 SSVKKS	265 5		275		285	290	295	300

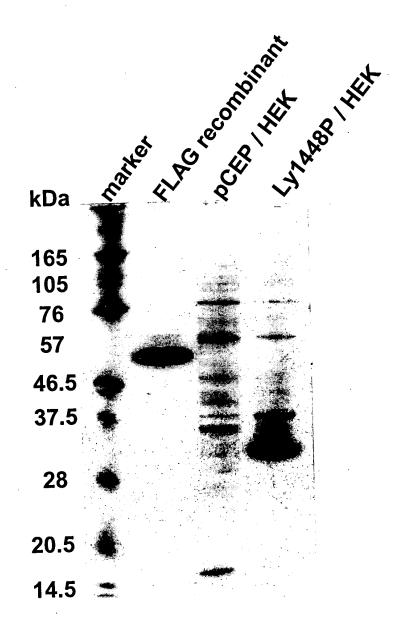


FIG. 8

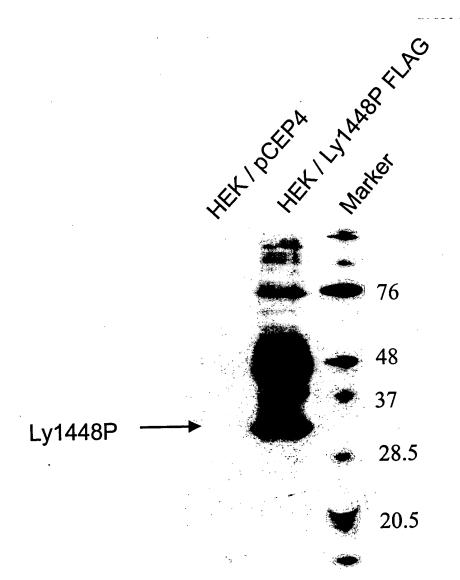


FIG. 9